(i) a 15 to 26-residue deleted peptide or deleted peptide analogue comprising formula (I) which forms an amphipathic  $\alpha$ -helix in the presence of lipids and in which one or two helical turns of the peptide or peptide analogue are optionally deleted:

$$Z_{1} - X_{1} - X_{2} - X_{3} - X_{4} - X_{5} - X_{6} - X_{10} - X_{10} - X_{10} - X_{11} - X_{12} - X_{13} - X_{14} - X_{15} - X_{16} - X_{17} - X_{18} - X_{19} - X_{20} - X_{21} - X_{22} - X_{23} - Z_{21} - X_{22} - X_{23} - Z_{22} - X_{23} - Z_{23} - Z_{24} - Z_{24} - Z_{25} -$$

or a pharmaceutically acceptable salt thereof, wherein:

- X<sub>1</sub> is Pro (P), Ala (A), Gly (G), Gln (Q), Asn (N), Asp (D) or D-Pro (p);
- X<sub>2</sub> is an aliphatic residue;
- $X_3$  is a Leu (L) or Phe (F);
- $X_4$  is Glu (E)
- X<sub>5</sub> is an aliphatic residue;
- $X_6$  is Leu (L) or Phe (F);
- $X_7$  is Glu (E) of Leu (L);
- $X_{g}$  is Asn (N) of Gln (Q);
- $X_0$  is Leu (L);
- $X_{10}$  is Leu (L), Trp (W) or Gly (G);
- $X_{11}$  is an acidic residue;
- $X_{12}$  is Arg (R);
- $X_{13}$  is Leu (L) or Gly (G);
- $X_{14}$  is Leu (L), Phe (F) or Gly (G);
- $X_{15}$  is Asp (D);
- $X_{16}$  is Ala (A);
- $X_{17}$  is Leu (L);
- $X_{18}$  is Asn (N) or Gln ( $\dot{Q}$ );
- $X_{19}$  is a basic residue;
- $X_{20}$  is a basic residue;
- $X_{21}$  is Leu (L);
- $X_{22}$  is a basic residue;
- X<sub>23</sub> is absent or a basic residue;
- $Z_1$  is  $R_2N$  or RC(O)NR-;
- $Z_2$  is -C (O) NRR or -C (O)  $\dot{Q}R$ ;

each R is independently -H,  $(C_1-C_6)$  alkyl,  $(C_1-C_6)$  alkenyl,  $(C_1-C_6)$  alkynyl,  $(C_5-C_{20})$  aryl,  $(C_6-C_{26})$  alkaryl, 5-20 membered heteroaryl or 6-26 membered alkheteroaryl or a 1 to 7-residue peptide or peptide analogue in which one more bonds between residues 1-7 are sindependently a substituted amide, an isostere of an amide or an amide mimetic;

each "-" between residues  $X_1$  to  $X_{23}$  and between residues of the peptide to  $Z_2$  independently designates an amide linkage, a substituted amide linkage, an isostere of an amide or an amide mimetic.

- 56. (Amended) The 15 to 26-residue deleted peptide or deleted peptide analogue of Claim 1, in which one helical turn is deleted.
- (Amended) The 15 to 26-residue deleted peptide or deleted peptide analogue of Claim 1, in which three, four, six, seven or eight residues  $X_1$ ,  $X_2$ ,  $X_3$ ,  $X_4$ ,  $X_5$ ,  $X_6$ ,  $X_7$ ,  $X_8$ ,  $X_9$ ,  $X_{10}$ ,  $X_{11}$ ,  $X_{12}$ ,  $X_{13}$ ,  $X_{14}$ ,  $X_{15}$ ,  $X_{16}$ ,  $X_{17}$ ,  $X_{18}$ ,  $X_{19}$ ,  $X_{20}$ ,  $X_{21}$  and  $X_{22}$  are deleted.
- 58. (Amended) The 15 to 26-residue deleted peptide or deleted peptide analogue of Claim 57, in which 3 consecutive residues are deleted.

59. (Amended) The 15 to 26-residue deleted peptide or deleted peptide analogue of Claim 57, in which 4 consecutive residues are deleted.

- 60. (Amended) The 15 to 26-residue deleted peptide or deleted peptide analogue of Claim 57, in which two non-contiguous sets of 3 consecutive residues are deleted.
- 61. (Amended) The 15 to 26-residue deleted peptide or deleted peptide analogue of Claim 57, in which two non-contiguous sets of 4 consecutive residues are deleted.
- 62. (Amended) The 15 to 26-residue deleted peptide or deleted peptide analogue of Claim 57, in which one set of 3 consecutive residues and one set of 4 consecutive residues are deleted.

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- 63. (Amended) The 15 to 26-residue deleted peptide or deleted peptide analogue of Claim 57, in which 6, 7 or 8 consecutive residues are deleted.
- 64. (Amended) The 15 to 26-residue deleted peptide or deleted peptide analogue of Claim 57, in which residues 19, 20 and 22 are not deleted.
- 65. (Amended) The 15 to 26-residue deleted peptide or deleted peptide analogue of Claim 57, in which residues 3, 6, 9 and 10 are not deleted.
- 66. (Amended) The 15 to 26-residue deleted peptide or deleted peptide analogue of Claim 1, in which  $X_{23}$  is absent.
- 67. (Amended) The 15 to 26-residue deleted peptide or deleted peptide analogue of Claim 1 in which: the "-" between residues designates -C (O) NH-;  $Z_1 \text{ is } H_2 \text{N-}; \text{ and}$   $Z_2 \text{ is -C (O) OH or a salt thereof.}$

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- (Amended) The 15 to 26-residue deleted peptide or deleted peptide analogue of Claim 1, in which the mean hydrophobic moment,  $\langle \mu_H \rangle$ , is about 0.45 to about 0.65.
- 69. (Amended) The 15 to 26-residue deleted peptide or deleted peptide analogue of Claim 68, in which the mean hydrophobic moment,  $\langle \mu_H \rangle$ , is about 0.50 to about 0.60.
- 70. (Amended) The 15 to 26-residue deleted peptide or deleted peptide analogue of Claim 1, in which the mean hydrophobicity,  $\langle H_o \rangle$ , is about -0.050 to about -0.070.
- 71. (Amended) The 15 to 26-residue deleted peptide or deleted peptide analogue of Claim 1, in which the mean hydrophobicity,  $\langle H_o \rangle$ , is about -0.030 to about -0.055.

- 72. (Amended) The 15 to 26-residue deleted peptide or deleted peptide analogue of Claim 1, in which the mean hydrophobicity of the hydrophobic face,  $\langle H_o^{pho} \rangle$ , is about 0.90 to about 1.20.
- 73. (Amended) The 15 to 26-residue deleted peptide or deleted peptide analogue of Claim 72, in which the mean hydrophobicity of the hydrophobic face, <H<sub>o</sub><sup>pho</sup>>, is about 0.94 to about 1.10.
- 74. (Amended) The 15 to 26-residue deleted peptide or deleted peptide analogue of Claim 1, in which the pho angle is about 160° to about 220°.
  - 75. (Amended) The 15 to 26-residue deleted peptide or peptide analogue of Claim 74, in which the pho angle is about 180° to about 200°.
- (Amended) A pharmaceutical composition comprising an ApoA-I agonist compound and a pharmaceutically acceptable carrier, excipient or diluent, wherein the ApoA-I agonist compound is a deleted peptide or deleted peptide analogue according to Claim 1 or 57.
  - 82. (Amended) The pharmaceutical composition of Claim 79 which is a lyophilized powder.

83. (Amended) The pharmaceutical composition of Claim 79 which is a solution.

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